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GAS NATURE EFFECT ON THE DESTRUCTION OF VARIOUS MICROORGANISMS UNDER CAVITATION ACTION

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Abstract. The disinfection of water from *Diplococcus*, *S.lutea*, *B.cereus*, and *Ps.fluorescens* bacteria types and the yeast *Sacch.cerevisiae via* the joint action of gas/US has been investigated. The investigated microorganisms were identified in natural water and wastewater in dominant amounts. The relative ranges of effective cell destruction by combined gas/US action have been established. The most efficient microorganism destruction has been obtained using Ar/US compared to He/US, CO₂/US and O₂/US and is accounted for by those gas properties able to affect sonochemical activity.

Keywords: cavitation, gas, water, microorganism, destruction.

1. Introduction

One of the most important tasks in water treatment technology is the development of methods which exclude chemical reagent usage such as chlorine or ozone, for the latter lead to the formation of toxic compounds [1, 2]. Recently researchers have turned their attention to physical methods of water treatment, especially the use of acoustic cavitation via ultrasound (US). Water disinfection under the influence of cavitation is supported by many research studies [3-8]. Acoustic cavitation is the phenomenon by which intense ultrasonic waves induce the formation, growth and subsequent collapse of dynamic microbubbles in a liquid [9].

The actions of cavitation alone, of bubbling gas alone and of the simultaneous action of cavitation and gas bubbling upon the disinfection of sewage water, natural water and microbial dispersions were considered in our previous study [10-12]. We observed that sonication enhanced the breakup of floc caused by microorganisms clumping together, which made them more susceptible to disinfection [11]. A mathematical model for the gas (Ar, He, O_2) effect on the process of water purification from *Bacillus* bacteria type in the absence of cavitation was proposed [13]. Our investigations showed that the

efficiency of microorganism destruction under the influence of cavitation was explained by cell membrane destruction induced by cavitation. A considerable number of studies were concentrated on the influence of cavitation on *Escherichia coli* cells, as traditionally they are the testorganisms used to evaluate the microbiological decontamination of water [1, 4-6, 8]. The effect of cavitation on the removal of fungi, *Enterococcus avium* and algae of *Microcystis aeruginosa, cyanobacteria* has also been presented [8, 14-17]. However, information on the cavitation effect on a wider range of various microorganisms is limited. The effect of cavitation is used not only for technological purposes, but also in the chemical industry, in particular in the process of modifying road bitumen [18], oxidation processes [19], *etc*.

The purpose of current investigation is to study the action of cavitation in the presence of different bubbled gases (Ar, He, CO₂, O₂) on the viability of *Diplococcus*, *Sarcina lutea*, *Bacillus cereus*, *Pseudomonas fluorescens* bacteria and the yeast (*Saccharomyces cerevisiae*) in aqueous media. The study of the different gases effect in a cavitation field, as well as the study of the behavior of a given gas in a cavitation field may help to determine the most efficient disinfection conditions for a range of microorganisms with different structural features.

2. Experimental

The investigated aqueous media were model dispersions prepared from natural water with added pure microorganism monocultures (*Diplococcus*, *Sarcina lutea*, *Bacillus cereus*, *Pseudomonas fluorescens* bacteria types and the *Saccharomyces cerevisiae* yeast). Thereby, five model dispersions containing different microorganisms were investigated.

The pure microorganism cultures were grown in test tubes in laboratory conditions at 310 K for 48 h (for bacterial cells) and at 303 K for 96 h (for yeast cells), on an agar nutrient medium with subsequent storage at 277 K. The agar nutrient medium composition is given below.

In our opinion, to simulate water disinfection under acoustic cavitation conditions, the microorganisms used in such studies need to be those most typically found in

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natural and sewage water. For this purpose, identification of microorganisms in different water sources was carried out (Table 1). So, in this study we investigated the effect of gas/US action on microorganisms that were previously identified in variously contaminated water sources. The qualitative microorganism composition of these water sources is also presented in Table 1.

Microorganisms in water sources under investigation were investigated by microscopy (coloration by Gram, cell sizes) to determine cell morphology. Their physiological (response to oxygen after being injected in a solid substrate in a test tube and response during growth in a nutrient medium) and culture features (the nature of colony growth on a nutrient medium in Petri dishes: colony form and profile, size, color, surface, shine and transparency, structure and consistency, edges; colony growth characteristics on the cross-hatched surface of test tube substrate: growth intensity, cross hatch features, color and surface) were investigated as well (Tables 2 and 3).

"Crushed drop" preparations with Lugol solution as a coloring agent and fixed cells preparations with safranin as a coloring agent were used to study the morphological features of the cells. "Crushed drop" preparation involved the evaluation of water samples with the aid of a microscope.

The source of cavitation was an ultrasonic wave source, namely UZDN-2T generator operating at the frequency of 22 kHz generating 35 W. Ultrasonic vibrations were transmitted by the magnetostriction radiator, which was immersed in the water being studied. During the investigations, the glass reactor vessel containing the dispersions ($V = 75 \text{ cm}^3$) was cooled throughout the process with tap water.

All experiments were carried out at $T = 298\pm1$ K, P = 0.1 MPa, and process time t = 2 h. Our previous studies showed that in the absence of US/gas, microorganisms quantity did not vary over a two-hour period in all water sources under investigation.

The degree of microorganism destruction (D_d) after gas/US treatment was found to be equal to the number of microorganisms (NM) at any given time relative to the initial number of microorganisms (NM_0) expressed as a percentage.

The effects of various gases such as oxygen, argon, helium, and carbon dioxide were comparatively investigated. Throughout the entire process gases were bubbled at the rate of ~1 cm³/s into the water under investigation. Gas bubbling and sonication were applied simultaneously to the glass reactor vessel.

Samples of sonicated water (1 cm³) were poured onto the nutrient medium in Petri dishes: meat water (1 dm³), peptone (10 g), agar (15 g) for bacteria and a malty mash (1 dm³) containing solids (6-8 %) and agar (2 %) for the yeast. Petri dishes were placed in a TS-80M-3

thermostat at $310\,\mathrm{K}$ for $48\,\mathrm{h}$ (for bacterial cells) and at $303\,\mathrm{K}$ for $96\,\mathrm{h}$ (for yeast cells). Water samples withdrawn before cavitation action were used as reference ones. The concentration of microbial cells was determined by the number of colony forming units (CFU), assuming that each colony developed from a single cell.

3. Results and Discussion

In order the qualitative content of the model microorganism dispersions is approached the actual content of microorganisms in contaminated waters, we investigated natural and waste waters (Table 1). Thus, bacteria *Sarcina, Diplococcus, Pseudomonas, Bacillus* and the yeast *Saccharomyces cerevisiae* were revealed and subjected to this study. The difference in their morphological, physiological [20] and culture [21] features is significant during the analysis of cell destruction in an aqueous medium (Tables 2 and 3).

Colony culture and cell morphology features are shown in Fig. 1: colony growth characteristics in Petri dishes on a nutrient medium (Ia-Va); fixed bacterial cells preparation with safranin as a coloring agent (Ib-IVb) and "crushed drop" preparation with Lugol solution as a coloring agent for yeast cells (Vb).

Natural water No1 was used for further investigations, as much as the simultaneous action of gas/US on the disinfection of waste and natural waters was presented in our previous work [13].

Depending on aqueous medium conditions and water sampling, the NM dynamics was established based upon a wide range of microbiological studies. Natural water No1 was sampled periodically during June and July. Some insignificant NM variations were observed for 10 days, within the narrow range from 820 to 1080 CFU/cm³, under relatively stable weather conditions, $T_{\rm H2O} = 286\pm2$ K. A sharp increase in water source temperature to 294 ±1 K caused a rapid increase of NM (up to 2090 CFU/cm³) for a short time, but over next two weeks there was practically no change in NM (Fig. 2).

As a result, natural water No1, with minimum NM_0 (820 CFU/cm³), was additionally infected by *Diplococcus, S.lutea, B. cereus, Ps. fluorescens* bacteria types and the *Sacch. cerevisiae* yeast, *i.e.* an individual model dispersion was created for each microorganism type. The results of gas/US action on these dispersions are given below.

The experimental results about the effect of acoustic destruction upon microorganisms increase the possibility of water disinfection by gas/US and the possibility of establishing the most effective conditions under which this process will lead to a high degree of microorganism destruction.

Microorganisms detected in the water objects

Investigated water objects	NM_0 , CFU/cm ³	Identified microorganisms
Natural water No1	820-2090	Bacterium, Sarcina, Bacillus* (35 %), Pseudomonas
Natural water No2	2200-2340	Staphylococcus, Pseudomonas, Bacillus* (40 %)
Natural water No3	1800-2050	Micrococcus, Sarcina Streptococcus, Bacterium* (29 %), Bacillus* (32 %)
Wastewater No1	110000	Micrococcus, Sarcina, Bacterium, Pseudomonas Bacillus* (25 %), Saccharomyces
Wastewater No2	38600	Micrococcus, Diplococcus, Streptococcus, Sarcina, Bacterium, Bacillus* (22 %)

Note: * dominant amounts.

Table 2

Morphological and physiological features of microorganisms

	Morphological features			Physiological features			
Investigated	Coloration by Gram	Sizes		Response to oxygen (the intensity of growth)			
microobjects		Width,	Length,	After being injected in a	After being injected in a melted substrate in a		
		μт μт		solid substrate in a test tube	test tube		
Bacillus cereus	Gram +	1.3	3.42	The superficial thin layer was created. Superficial layer: all-round, dense, wrinkled, plicate. Isolated colonies in the deep of medium along the injection were revealed.	Evenly distributed superficial thin layer of colonies was found. Superficial layer is the same as in the case of solid medium. Growth of single colonies over the medium and distributed on the wall of the tube was revealed. Moderate growth of colonies along the depth of medium in vitro was revealed. Evenly distributed superficial thin layer was investigated. Superficial layer is the same as in the case of solid medium. Explicit growth of colonies distributed on the tube wall over the medium.		
Diplococcus	Gram +	0.6	-	The superficial thin layer was present. Superficial layer: all-round, dense, smooth.			
Sarcina lutea	Gram +	2.2	_	The same	The same		
Pseudomonas fluorescens	Gram –	0.53	1	Surface growth of the colonies and intensive colonies growth along the injected medium in vitro was shown. Superficial layer: all-round, dense, smooth.	The superficial thin layer was evenly distributed on the medium surface. Superficial layer is the same as in the case of solid medium. Growth of single colonies on the wall of the tube over the environment was found. Intensive colonies growth throughout the medium depth in vitro was present.		
Saccharomyces cerevisiae	Gram +	5.17	8.67	Superficial layer: discontinuous, but thick and dense. Convex colonies growing upwards.	The superficial thin layer was unevenly distributed and relatively thick. Growth of single colonies above medium was a little distributed on the wall of the tube.		

Table 3

Cultural features of microorganisms

Investigated microobjects	Characteristics of colony growth on a nutrient medium in Petri dishes							
	Shape and profile	Size, mm	Color	Surface	Shine and transparency	Structure and consistency	Edges	
1	2	3	4	5	6	7	8	
Bacillus cereus	compact, fold, hilly; curved profile	<1; 1–4	white	wrinkled with folded center	opaque, matt, moisture	finely granular (farinaceous); mucous texture	wavy, winding	
Diplococcus	compact, round; convex or flat profile	<1; 1–3	peachy; produce orange pigment	smooth	opaque, brilliant, moisture	homogeneous; soft texture	smooth, flat	

Table 3 (continued)

1	2	3	4	5	6	7	8
Sarcina lutea	compact, round; convex or flat profile	<1; 1–3	citric; produce yellow pigment	smooth	opaque, brilliant, moisture	homogeneous; soft texture	smooth, flat
Pseudomonas fluorescens	compact, round; convex or flat profile	<1; 1–2	grayish (dirty white) or without color; produce green- yellow fluorescent pigment	smooth	opaque, brilliant, moisture	homogeneous; soft texture	smooth, flat
Saccharomyces cerevisiae	round; colonies in the deep have cross-like borders, which burst an agar; curved profile	1–5	gray-white	hilly or slightly wrinkled	opaque, brilliant, sometimes matt	homogeneous; texture is oily (pasty) and thick	flat, sometimes wavy

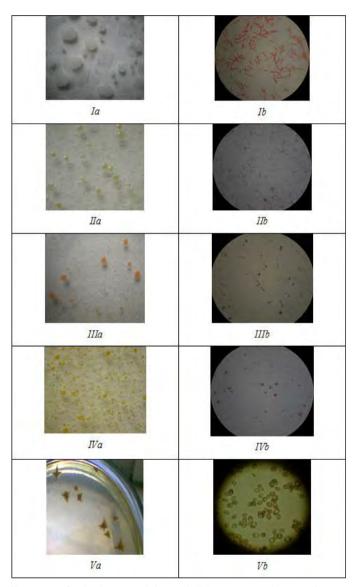


Fig. 1. Character of colonies growth in Petri dishes on a nutrient medium (a) and SEM images of microorganism cells (b) for *Bacillus cereus* (I); *Sarcina lutea* (II); *Diplococcus* (III); *Pseudomonas fluorescens* (IV) and *Saccharomyces cerevisiae* (V). Magnification of 1200× (I); 1500× (II-IV) and 480× (V)

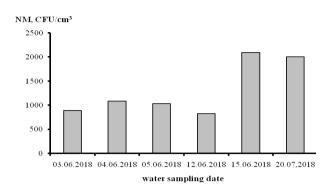


Fig. 2. Dynamics of NM_0 changes in the natural water No1 depending on the sampling date (sampling depth was 100 mm from the water surface)

3.1. Destruction of *Bacillus Cereus*Bacteria Type

The effect of simultaneous gas/US action upon bacillary spore formation in the *Bacillus cereus* bacterium (Fig. 1, Ib) is shown in Fig. 3.

Consequently, one may compare *Bacillus* destruction within a cavitational field for different gases being bubbled into its dispersion and so determine the gaseous atmosphere in which this process is most effective: $Ar/US > O_2/US > He/US > CO_2/US$

It has been established that the most effective destruction process for the *Bacillus* bacteria occurs in the presence of argon in a cavitational field.

3.2. Destruction of *Pseudomonas Fluorescens* Bacteria Types

Because bacillary *Ps. fluorescens* type bacteria (Fig. 1, IIb), unlike *Bacillus* microbial cells, are unable to form spores, they were investigated for their response to gas/US treatment.

The NM_0 values in microbial dispersions were: $9.5 \cdot 10^4$; $9.7 \cdot 10^3$ and $1.8 \cdot 10^3$ CFU/cm³ for O₂/US, CO₂/US and Ar/US, respectively. The process of water purification, which is subjected to a gas/US process over a two-hour period, is shown in Fig. 4.

It should be noted that during the first stage of being subjected to He/US effect (t = 1800 s) an increase in NM was observed as a result of intensive disaggregation of microbial cells found in the effluent water, whilst active microorganism destruction ($t_{\text{He/US}} = 1800-7200 \text{ s}$) was only observed at the second stage. Therefore, to determine D_d values, the value of NM_{max} was considered instead of NM_0 . For this reason, in this particular instance the comparison between the effect of helium and other gases in a cavitational field is inappropriate (this is not shown in the figure).

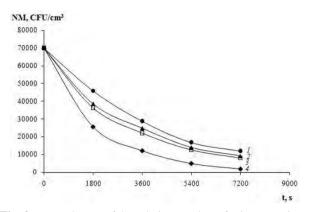


Fig. 3. Dependences of the relative number of microorganisms (*Bacillus* bacteria type) on the process time under the gas/US action: CO_2/US (1), He/US (2), O_2/US (3), Ar/US (4). Conditions: $NM_0 = 7 \cdot 10^4$ CFU/cm³, $T = 298 \pm 1$ K

The effective destruction order for *Ps. fluorescens* bacteria types for different gases being bubbled in a cavitational field is as follows:

 $Ar/US > CO_2/US > O_2/US > He/US$

The most effective rate constants for the destruction of these bacteria (3.96±0.02·10⁻⁴ s⁻¹) was established for Ar/US action.

3.3. Destruction of *Diplococcus* and *Sarcina Lutea* Bacteria Types

Although *Diplococcus* and *Sarcina lutea* bacteria types from the *Coccaceae* family are related to spherical microorganisms (Fig. 1, IIIb and Fig.1, IVb, respectively), they differ by size [22] and by numbers of connected cells (*Diplococcus* cells are connected in pairs, whilst *Sarcina* cells are in blocks).

The general effect of gas/US action on bacteria viability is shown in Fig. 5. According to calculated effective rate constants for microorganisms destruction, both Diplococcus ($k_d = 3.29 \pm 0.03 \cdot 10^{-4} \, \mathrm{s}^{-1}$) and S.lutea ($k_d = 5.95 \pm 0.07 \cdot 10^{-4} \, \mathrm{s}^{-1}$) are destroyed most readily in the presence of argon rather than in helium. The value of k_d is less by 0.52 and 0.55 times under He/US for Diplococcus and S.lutea, respectively.

Comparing the efficiency of the process for monocultures, more active destruction of *Sarcina* (both in argon and in helium) is observed, which may be related to its significantly larger cell size compared to that of *Diplococci* (\sim 3.5 times) [22]. Thus, the destruction of *Diplococcus* and *S.lutea* under cavitation conditions has been shown to occur most effectively in argon. D_d is approximately twice higher under Ar/US condition than for He/US as shown by the curves of the relative number of microorganisms in the figure. The influence of oxygen and carbon dioxide under cavitation conditions is not presented in the figure, since under these conditions processes of disaggregation of microbial cell clusters were observed and such processes were described in [23].

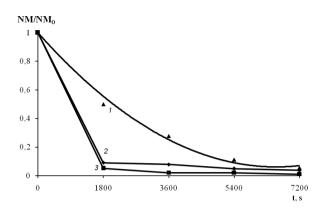


Fig. 4. Dependences of the relative number of microorganisms (*Ps.fluorescens* bacteria type) on the process time under the gas/US action: O_2 /US (1), CO_2 /US (2), Ar/US (3). Conditions: $NM_0 = 9.5 \cdot 10^4$ for O_2 /US; $NM_0 = 9.7 \cdot 10^3$ for CO_2 /US; $NM_0 = 1.8 \cdot 10^3$ for Ar/US; $T = 298 \pm 1$ K

3.4. Destruction of the Saccharomyces Cerevisiae Yeast

According to Fig. 6, the efficiency of *Sacch. cerevisiae* destruction (Fig. 1, Vb) when subjected to inert gas bubbling for $t_{\rm gas/US} = 5400-7200$ s remains practically unchanged, compared to that for oxygen and carbon dioxide. The yeast suspension was >98 % after one hour of the process duration, which confirms the previous results [14]. The number of yeast cells per 1cm³ did not exceed 100 cells after $t_{\rm gas/US} = 3600$ s. This mean that in accordance with acceptable microbiological parameters the investigated water meets fully with drinking water requirements.

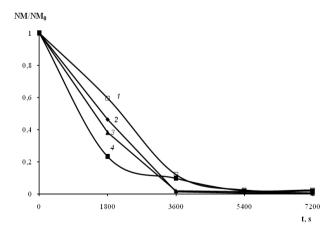


Fig. 6. Dependences of the relative number of yeasts (*Sacch. cerevisiae*) on the process time under the gas/US action: He/US (1), O₂/US (2), Ar/US (3), CO₂/US (4). Conditions: $NM_0 = (4.2-5.2) \cdot 10^3 \text{ CFU/cm}^3$ for He/US and CO₂/US; NM₀= (1.9–2.07) · 10⁴ CFU/cm³ for O₂/US and Ar/US; $T = 298 \pm 1 \text{ K}$

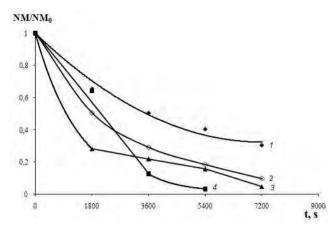


Fig. 5. Dependences of the relative number of microorganisms (*Diplococcus* (1, 2) and *S.lutea* (3, 4) bacteria types) on the process time under the gas/US action: He/US (1, 3) and Ar/US (2, 4). Conditions: $NM_0 = 1.1 \cdot 10^5$ CFU/cm³ (1, 2) and $NM_0 = 6.4 \cdot 10^3$ CFU/cm³ (3, 4); $T = 298 \pm 1$ K

The destruction effectiveness order for *Sacch*. *cerevisiae* when subjected to various gas/US combinations is as follows: $Ar/US > He/US > CO_2/US > O_2/US$.

As a result, the yeast suspension was almost entirely disinfected under cavitation conditions for various bubbled gases in a relatively short time period compared to that for microbial dispersions. It should be noted that the unicellular yeast type fungi studied (eukaryotes) differ significantly in their morphology from the prokaryotes in this study. Yeast cells, unlike bacterial ones, although characterized by stronger membranes, are almost 10 times larger [22]. This may have a direct bearing upon the destruction effect experienced under cavitation conditions.

4. Conclusions

This study has yielded the following results:

The destruction of *Diplococcus*, *S.lutea*, *B.cereus*, *Ps.fluorescens* bacteria types and the *Sacch.cerevisiae* yeast under cavitation action depends on the nature of the bubbled gas. The relative destruction efficiency for cells subjected to gas/US shows the greatest water disinfection efficiency for argon compared to helium, oxygen and carbon dioxide.

Regardless of the types of microorganism studied, their morphology, and their concentration per unit volume of dispersion, the highest effectiveness of microorganism destruction in the aqueous medium was observed under the simultaneous action of argon and cavitation.

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ВСТАНОВЛЕННЯ ЕФЕКТИВНОЇ ПРИРОДИ ГАЗУ НА РУЙНУВАННЯ РІЗНИХ МІКРООРГАНІЗМІВ ПІД ДІЄЮ КАВІТАЦІЇ

Анотація. Досліджено знезараження води від бактерій роду Diplococcus, S.lutea, B.cereus, Ps.fluorescens та дріжджів Sacch.cerevisiae за одночасної дії газ/V3 як мікроорганізми, які були визначені в природних та стічних водах в домінуючій кількості. Встановлені відносні ряди ефективного руйнування клітин під дією газ/V3. Найбільша ефективність знищення мікроорганізмів було досягнуто під дією Ar/V3, порівняно з He/V3, $CO_2/V3$ і $O_2/V3$ що пояснюється властивостями газу, здатними впливати на сонохімічну активність.

Ключові слова: кавітація, газ, вода, мікроорганізми, руйнування.